

Ecosystem Services and Trade-Offs Mediated by Urban Water Bodies for Homeless Populations in Phoenix

Wolf, A.^{1,2}, M.M. Palta,² N.B. Grimm,² J. Gwiszcz,³ O. Schwake,⁴

¹Arizona State University Undergraduate Program; ²School of Life Sciences, ASU; ³School of Human Evolution and Social Change, ASU; ⁴School of Sustainable Engineering and the Built Environment, ASU



INTRODUCTION

- Urban runoff entering stormwater outfalls in Phoenix have created “accidental” wetlands in the otherwise dry Salt River bed
- Urban wetland environments provide heat mitigation, privacy, and other services to the homeless population in Phoenix but high pathogen loading at outfalls presents public health concerns

STUDY OBJECTIVES

- Assess the extent of use and the ecosystem services and trade-offs provided by the wetlands to the homeless population in Phoenix
- Monitor *Escherichia coli* (*E. coli*) loading levels and wetland-mediated attenuation rates of *E. coli* at points along flowpaths from six outfalls
- Determine source of fecal contamination at outfalls by analyzing environmental *E. coli* isolates’ protein profiles with *E. coli* strains of human origin

SAMPLING DESIGN

- Water samples collected during baseflow and storm conditions May–July 2013
- Samples collected at 3–5 points along flowpaths from outfalls and through wetland areas



★ = Sampling location

PHYSICAL VARIABLES IN OUTFALLS DURING BASE AND STORMFLOW

- Measured discharge, flow rate, dissolved oxygen (DO), and conductivity

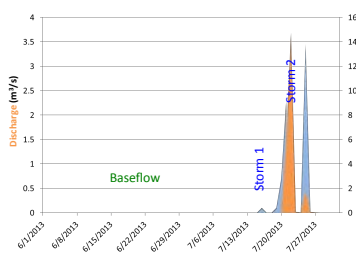
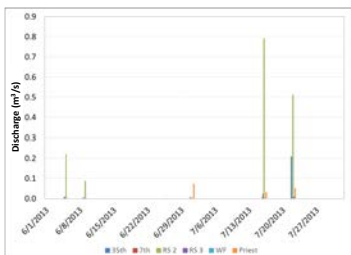


Figure 1: (a) Precipitation record and discharge exiting Priest site (USGS gage) during study period; (b) discharge measured in study outfalls on collection days.



PATHOGEN LOADING AND ATTENUATION

- Water samples collected at sampling points analyzed within 12 hrs for coliform and *E. coli* levels by plating 500 μ L on selective media (Brilliance®) and incubating at 70 °C for 20 hrs
- Attenuation along flowpath quantified as disappearance of *E. coli* colonies per 1 mL between loading at outfall and end of flowpath

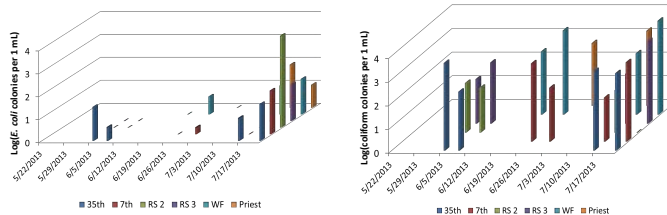


Figure 2: *E. coli* (left) and coliform (right) loading at outfalls. Coliform levels remained consistently high during base and storm flow; *E. coli* levels spiked during storms.

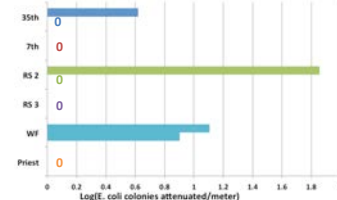


Figure 3: Attenuation rate of *E. coli* colonies between beginning (at outfall) and end of flowpaths during storm events. Little to no attenuation was observed during baseflow conditions.

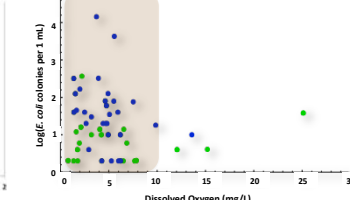


Figure 4: Baseflow (green) and storm (blue) *E. coli* concentrations at all sites regressed against dissolved oxygen; the data suggest a decrease in *E. coli* vitality below 10 mg/L O₂. Temperature, conductivity, and flow rate did not demonstrate a relationship with *E. coli* concentration.

CONCLUSIONS

- Pathogen and *E. coli* loading at outfalls well exceed bathing and drinking water standards; levels also peak under storm conditions
- Wetland-mediated attenuation of *E. coli* from outfall to wetland area (end of flowpath) was observed under storm conditions
- Potential DO cut-off for environmental *E. coli* exists near 10 mg/L
- Preliminary MALDI-TOF-MS results show strain-level clustering of environmental with human strain *E. coli* isolates

FUTURE DIRECTION

- Follow up on trash monitoring
- Nutrient analysis and analysis of other environmental variables along flowpaths may suggest ways in which wetland environments may remove pathogens
- Sucralose testing to support fecal contamination is of human origin

SOURCE TRACKING

- Approx. 120 *E. coli* colonies were isolated from Brilliance® media and purified by twice-culturing on TSA media and storing at -80 °C
 - E. coli* of known human origin: raw sewage samples collected from Scottsdale Water Campus (SCW); plated within 36 hrs, cultured over 20 hrs
 - E. coli* from environmental samples: samples collected at study sites; plated within 12 hrs, cultured over 20 hrs
- Protein extractions of isolated colonies were prepared with formic acid and Acetonitrile with alpha-Cyano-4-hydroxycinnamic acid matrix solution
- Protein profiles of environmental and human strain *E. coli* isolates analyzed using MALDI-TOF-MS

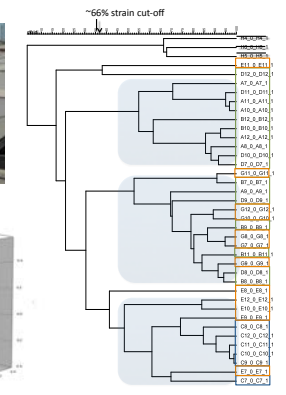
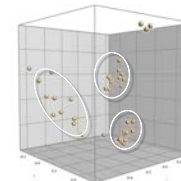


Figure 5: Select MALDI-TOF MS results. Right: Strain-level dendrogram; Orange = human derived (E: SCW strain #1; G: SCW strain #3), Green = baseflow (A: 35th Ave outfall; B: RS Central Ave outfall #2; D: Priest outfall), Blue = storm flow (C: RS 7th Ave outfall). Apparent clustering of *E. coli* strains from 35th Ave & Priest, RS Central Ave #2 & SCW strain #1, and RS 7th Ave & SCW strain #3. Left: 3D visualization of clustering.

ACKNOWLEDGEMENTS

Thank you to the members of Nancy Grimm’s lab (Jeremiah McGehee, Marena Sampson, Truman Combs, Erin Worth, Amalia Handler, & Emma Holland) for assistance in the field and in the lab; and a special thank you to Morteza Abbaszadegan and Todd Sandrin and the members of their labs for providing their facilities and expertise. Funding was provided by CAP LTER